

SEQUENCE LISTING

<110> The Australian National University

5 <120> METHOD OF PRODUCING PLANTS HAVING ENHANCED TRANSPIRATION EFFICIENCY AND  
PLANTS PRODUCED THEREFROM

<130> 94948/MRO

10 <150> AU PS3339  
<151> 2002-07-02

<160> 45

15 <170> PatentIn version 3.1

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Asp Gly Glu Ile Ser Pro Ala Ile Gly Asp Leu Lys Ser Leu Leu Ser  
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5 Ser Gln Ile Pro Asn Leu Lys Ile Leu Asp Leu Ala Gln Asn Lys Leu  
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Ser Gly Glu Ile Pro Arg Leu Ile Tyr Trp Asn Glu Val Leu Gln Tyr  
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Gly Phe Leu Gln Val Ala Thr Leu Ser Leu Gln Gly Asn Gln Leu Ser  
 30 260 265 270

Gly Lys Ile Pro Ser Val Ile Gly Leu Met Gln Ala Leu Ala Val Leu  
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40 Asn Leu Thr Phe Thr Glu Lys Leu Tyr Leu His Ser Asn Lys Leu Thr  
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665

670

Val Ala Ile Lys Arg Leu Tyr Ser His Asn Pro Gln Ser Met Lys Gln  
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Lys Asp Leu Gly Val Val Lys Lys Val Phe Gln Leu Ala Leu Leu Cys  
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265

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Thr Ile Pro Ala Glu Leu Gly Lys Leu Glu Glu Leu Phe Glu Leu Asn  
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Leu Ala Asn Asn Asn Leu Gln Gly Pro Ile Pro Ala Asn Ile Ser Ser  
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40 Leu Ser Ser Asn Asn Phe Lys Gly Asn Ile Pro Ser Glu Leu Gly His  
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Ile Ile Asn Leu Asp Thr Leu Asp Leu Ser Tyr Asn Glu Phe Ser Gly  
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5 Pro Val Pro Ala Thr Ile Gly Asp Leu Glu His Leu Leu Glu Leu Asn  
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Leu Ser Lys Asn His Leu Asp Gly Pro Val Pro Ala Glu Phe Gly Asn  
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Leu His His Asp Cys Asn Pro Arg Ile Ile His Arg Asp Val Lys Ser  
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Gln Leu Ile Leu Ser Lys Ala Asp Asp Asn Thr Val Met Glu Ala Val  
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Phe Cys Asp Asn Val Ser Tyr Ser Val Val Ser Leu Asn Leu Ser Ser  
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Ala Thr Leu Thr Gln Ile Pro Asn Leu Lys Arg Leu Asp Leu Ala Gly  
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Leu Gln Tyr Leu Gly Leu Arg Gly Asn Met Leu Thr Gly Thr Leu Ser  
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Ser Asp Met Cys Gln Leu Thr Gly Leu Trp Tyr Phe Asp Val Arg Gly  
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Arg Leu Thr Gly Arg Ile Pro Glu Val Ile Gly Leu Met Gln Ala Leu  
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30 Met Leu Thr Gly Pro Ile Pro Ser Glu Leu Gly Asn Met Ser Arg Leu  
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Ser Tyr Leu Gln Leu Asn Asp Asn Lys Leu Val Gly Thr Ile Pro Pro  
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Glu Leu Gly Lys Leu Glu Gln Leu Phe Glu Leu Asn Leu Ala Asn Asn  
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Thr Leu Gly Asp Leu Glu His Leu Leu Ile Leu Asn Leu Ser Arg Asn  
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His Leu Ser Gly Gln Leu Pro Ala Glu Phe Gly Asn Leu Arg Ser Ile  
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Gln Met Ile Asp Val Ser Phe Asn Leu Leu Ser Gly Val Ile Pro Thr  
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Glu Leu Gly Gln Leu Gln Asn Leu Asn Ser Leu Ile Leu Asn Asn Asn  
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	595	600	605
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Gln Lys Lys Ile Leu Gln Gly Ser Ser Lys Gln Ala Glu Gly Leu Thr			
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Ile Met Arg Val Thr Glu Asn Leu Asn Glu Lys Phe Ile Ile Gly Tyr			
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Gly Ala Ser Ser Thr Val Tyr Lys Cys Ala Leu Lys Ser Ser Arg Pro			
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Ile Ala Ile Lys Arg Leu Tyr Asn Gln Tyr Pro His Asn Leu Arg Glu			
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Phe Glu Thr Glu Leu Glu Thr Ile Gly Ser Ile Arg His Arg Asn Ile			
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Val Ser Leu His Gly Tyr Ala Leu Ser Pro Thr Gly Asn Leu Leu Phe			
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Tyr Asp Tyr Met Glu Asn Gly Ser Leu Trp Asp Leu Leu His Gly Ser  
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5 Leu Lys Lys Val Lys Leu Asp Trp Glu Thr Arg Leu Lys Ile Ala Val  
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Gly Ala Ala Gln Gly Leu Ala Tyr Leu His His Asp Cys Thr Pro Arg  
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Ile Ile His Arg Asp Ile Lys Ser Ser Asn Ile Leu Leu Asp Glu Asn  
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Thr Lys Arg Asn Pro Leu Glu Arg Pro Thr Met Leu Glu Val Ser Arg  
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Val Leu Leu Ser Leu Val Pro Ser Leu Gln Val Ala Lys Lys Leu Pro  
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10 Ser Leu Asp His Ser Thr Lys Lys Leu Gln Gln Glu Asn Glu Val Arg  
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Thr Glu Leu Glu Thr Val Gly Ser Ile Arg His Arg Asn Leu Val Ser  
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His Arg Asp Val Lys Ser Ser Asn Ile Leu Leu Asp Glu His Phe Glu  
 385 390 395 400

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Ala His Leu Ser Asp Phe Gly Ile Ala Lys Cys Val Pro Ala Ala Lys  
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 5 cgaacgtcta cgtcttcccc tacaccatgt tctgcgagat ggcctcggcg aacttgagga 240  
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 15 ggagcgccag ctggaacagc ttcttcacct cggccaggtc cttgcaggng tctcccacgt 540  
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 40 caccacctcg ggcac 486  
  
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<211> 509  
 <212> DNA  
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 10 cctgcgctgc gtccacāāā atgtcagtca aactactccc tgcaatccgc ctcaactcaag 180  
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 15 tggcccgca gctgacgtac tcgtcgacgt agggccggcg cgacggctgc ggcggcagct 360  
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25 <210> 26  
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 35 accctagcta ctgagtccca tgtaatctc ctgcgctgcg tccacāāāā tgtcagtcaā 180  
 actactcccc ctgcaatccg cctcaactcāā ggcgcctcac cgaacgtcta cgtcttcccc 240  
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 40 ctggācgaat tggccgag 318

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15

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40

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gcgtccacaca aaatgtcagt caaactactc cctgcaatcg gcctcatttt ttgtgtgtcc 180

tcaccgaacg tctacgtctt cccctacacc atgttctgcg agatggcctc gccgaacttg 240

5 aggaacagct cggcgtccga ggtgctggac gagttggcgc aggaaagggc gccggtgccc 300

cgcaggctga cgtactcgtc gacgtaggcc ggcggcgacg gctgcggcgg cagctggtgg 360

10 tgcgctgctg ccttctgctg ctgctgctgc ggctgcggcg gcggctccgg gttcaccagg 420

cagtcaaggc cgcgcaccac ctctgtcatc gtcggccggc ccgagggctg ccgcttggtg 480

cagaggagcg ccagctggaa cagcttcttc acctcgccca ggtccttgca ggtgtctccc 540

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<213> partial maize ERECTA

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tgctcttgcg tcacatgact ttttttacag ctaacaacac cctagctact gagtcccatg 180

30 ttaatctcct gcgctgcgtc ccacaa 206

<210> 31

35 <211> 534

<212> DNA

<213> partial maize ERECTA

<400> 31

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<213> partial maize ERECTA

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<210> 34

20 <211> 533

<212> DNA

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15

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20

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683

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25 gtacgtcagc ctgcggggca ccggcgccct ctctgcgcc aactcgtcca gcacctcgga 540
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10  
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35 Ser Ile Pro Pro Glu Leu Gly Arg Leu Thr Gly Leu Phe Asp Leu Asn  
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40 Leu Ala Asn Asn His Leu Glu Gly Pro Ile Pro Asp Asn Leu Ser Ser  
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Cys Val Asn Leu Asn Ser Phe Asn Ala Tyr Gly Asn Lys Leu Asn Gly  
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Thr Ile Pro Arg Ser Leu Arg Lys Leu Glu Ser Met Thr Tyr Leu Asn  
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